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# 8-[(2-Hydroxyphenyl)imino]-3,5a,9-trimethyl-3a,4,5,5a,8,9b-hexahydro-naphtho[1,2-b]furan-2(3H)-one

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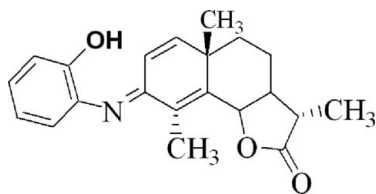
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Key indicators: single-crystal X-ray study;  $T = 273$  K; mean  $\sigma(\text{C}-\text{C}) = 0.005$  Å; R factor = 0.041; wR factor = 0.101; data-to-parameter ratio = 8.6.

The title compound,  $\text{C}_{21}\text{H}_{23}\text{NO}_3$ , is a phenylimine derivative of the well known anthelmintic agent  $\alpha$ -santonin. The *trans*-fused cyclohexane and  $\gamma$ -lactone rings of the  $\alpha$ -santonin ring system adopt chair and envelope conformations, respectively, whereas the hexadiene ring is approximately planar [maximum deviation =  $0.029(4)$  Å] and forms a dihedral angle of  $62.30(11)^\circ$  with the benzene ring. An intramolecular  $\text{O}-\text{H}\cdots\text{N}$  hydrogen bond is observed.

## Related literature

For the isolation and anthelmintic use of  $\alpha$ -santonin, see: Miana & Al-Lohedan (1986). For the crystal structure and stereochemistry of  $\alpha$ -santonin, see: White & Sim (1975); Coggon & Sim (1969). For the crystal structure of a related compound, see: Yousuf *et al.* (2012).



## Experimental

### Crystal data

 $\text{C}_{21}\text{H}_{23}\text{NO}_3$  $M_r = 337.40$ 

Orthorhombic,  $P2_12_12_1$   
 $a = 8.6000(9)$  Å  
 $b = 10.7458(11)$  Å  
 $c = 19.729(2)$  Å  
 $V = 1823.2(3)$  Å<sup>3</sup>

$Z = 4$   
 Mo  $K\alpha$  radiation  
 $\mu = 0.08$  mm<sup>-1</sup>  
 $T = 273$  K  
 $0.54 \times 0.14 \times 0.04$  mm

### Data collection

Bruker SMART APEX CCD area-detector diffractometer  
 Absorption correction: multi-scan (SADABS; Bruker, 2000)  
 $T_{\min} = 0.957$ ,  $T_{\max} = 0.997$

10874 measured reflections  
 1955 independent reflections  
 1385 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.055$

### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.041$   
 $wR(F^2) = 0.101$   
 $S = 1.04$   
 1955 reflections

228 parameters  
 H-atom parameters constrained  
 $\Delta\rho_{\max} = 0.14$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.13$  e Å<sup>-3</sup>

**Table 1**

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{O3}-\text{H3A}\cdots\text{N1}$	0.82	2.28	2.747 (4)	116

Data collection: SMART (Bruker, 2000); cell refinement: SAINT (Bruker, 2000); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: SHELXTL (Sheldrick, 2008); software used to prepare material for publication: SHELXTL, PARST (Nardelli, 1995) and PLATON (Spek, 2009).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: RZ2770).

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## supplementary materials

*Acta Cryst.* (2012). E68, o2158 [doi:10.1107/S1600536812027146]

## 8-[(2-Hydroxyphenyl)imino]-3,5a,9-trimethyl-3a,4,5,5a,8,9b-hexahydro-naphtho[1,2-*b*]furan-2(3*H*)-one

Sammer Yousuf, Syed M. Younas, Nida Ambreen, Khalid M. Khan and Ghulam A. Miana

### Comment

$\alpha$ -Santonin was isolated from *Artemisia santonica* (Miana & Al-Lohedan, 1986) and widely used in the past as an anthelmintic drug to expels parasitic worms (helminths) from the body, by either killing or stunning them. The title compound was prepared as a part of our ongoing reaserch to synthesise bioactive derivatives of  $\alpha$ -santonin *via* biology oriented synthesis (BIOS). The title compound is an analogue of our previously reported compound 3,5a,9-trimethyl-3a,5,5a,9b-tetrahydronaphtho[1,2-*b*]furan-2,8(3*H*,4*H*)-dione-8-(*N*-phenylhydrazone), with the difference that the phenylhydrazine moiety is replaced by a 2-hydroxyphenylimine group (C16–C21) attached to the  $\alpha$ -santonin ring system (O1–O2/C1–C15). The cyclohexadiene ring (C6–C11) is almost planar with a maximum deviation from the least square plane of 0.029 (3) Å for atom C7 and forms a dihedral angle of 62.30 (11)° with the phenyl ring. The cyclohexane ring (C3–C6/C11–C12) adopts a chair conformation [ $Q = 0.594$  (4) Å,  $\theta = 8.2$  (4)° and  $\varphi = 304$  (2)°] and is *trans* fused to the  $\gamma$ -lactone ring (O1/C1–C3/C12) which adopts an envelope conformation with atom C3 0.228 (3) Å out of the plane formed by the rest of the ring atoms. The two methyl substituents at atoms C6 and C2 exist in *axial* and *pseudo equatorial* orientations, respectively (Fig. 1). The bond dimensions are similar to those found in the structurally related compounds (Yousuf *et al.*, 2012; White & Sim, 1975; Coggon & Sim, 1969). An intramolecular O—H $\cdots$ N hydrogen bond is present (Table 1). In the crystal, molecules are arranged into layers parallel to the *ab* plane only by van der Waals forces (Fig. 2).

### Experimental

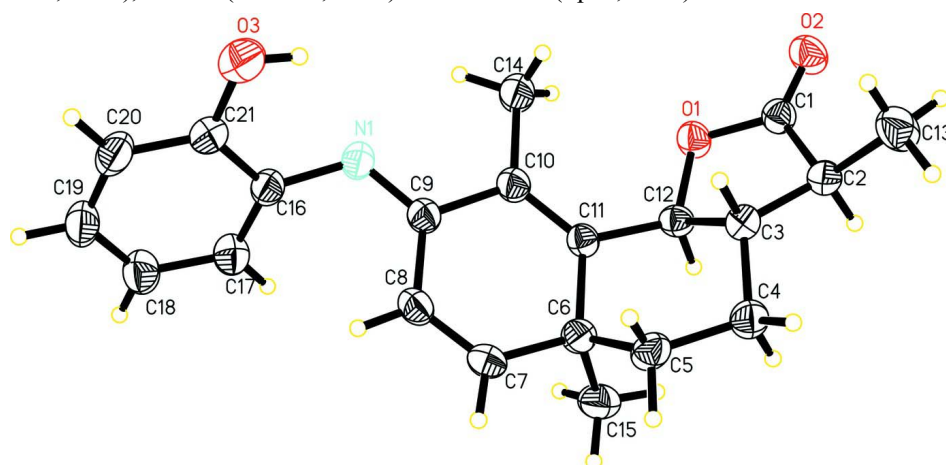
In a 100 ml round bottomed flask toluene (25 ml) and  $\alpha$ -santonin (400 mg, 1.6 mmol) were taken, then 2-amino phenol (11.2 mmol) was added with continuous stirring. The reaction mixture was refluxed and monitored by TLC. After completion of reaction (24 h), the mixture was cooled and extracted with water. The organic layer was dried over Na<sub>2</sub>SO<sub>4</sub>, filtered and the solvent evaporated under vacuum in a rotary evaporator. The crude product was chromatographed on a silica gel column using *n*-hexane:ethyl acetate (7:3 v/v) as mobile phase to obtain yellow crystals of title compound in 85% yield.

### Refinement

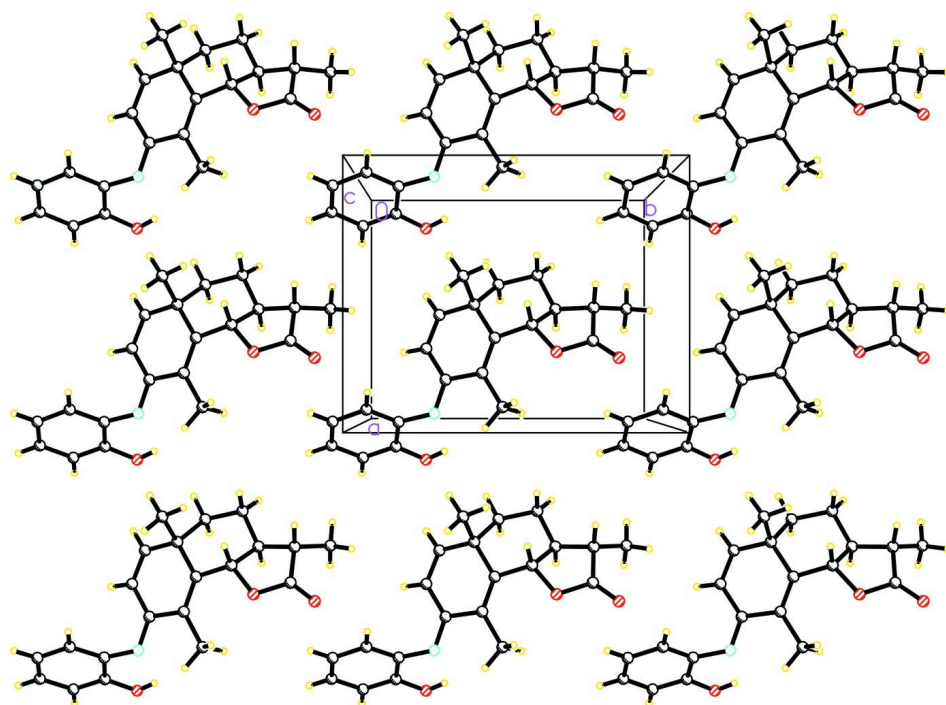
H atoms were positioned geometrically with C—H = 0.93–0.97 Å, O—H = 0.82 Å, and constrained to ride on their parent atoms with  $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{C})$  or  $1.5 U_{\text{eq}}(\text{C}, \text{O})$  for methyl and hydroxy H atoms. A rotating group model was applied to the methyl groups. 1433 Friedel pairs were merged.

**Computing details**

Data collection: *SMART* (Bruker, 2000); cell refinement: *SAINTE* (Bruker, 2000); data reduction: *SAINTE* (Bruker, 2000); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL* (Sheldrick, 2008), *PARST* (Nardelli, 1995) and *PLATON* (Spek, 2009).

**Figure 1**

The molecular structure of the title compound with displacement ellipsoids drawn at 30% probability level.

**Figure 2**

The crystal packing of the title compound viewed down the *c* axis.

8-[(2-Hydroxyphenyl)imino]-3,5a,9-trimethyl-3a,4,5,5a,8,9b-hexahydronaphtho[1,2-b]furan-2(3H)-one

Crystal data

$C_{21}H_{23}NO_3$	$F(000) = 720$
$M_r = 337.40$	$D_x = 1.229 \text{ Mg m}^{-3}$
Orthorhombic, $P2_12_12_1$	Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
Hall symbol: P 2ac 2ab	Cell parameters from 1151 reflections
$a = 8.6000 (9) \text{ \AA}$	$\theta = 2.8\text{--}18.6^\circ$
$b = 10.7458 (11) \text{ \AA}$	$\mu = 0.08 \text{ mm}^{-1}$
$c = 19.729 (2) \text{ \AA}$	$T = 273 \text{ K}$
$V = 1823.2 (3) \text{ \AA}^3$	Plate, yellow
$Z = 4$	$0.54 \times 0.14 \times 0.04 \text{ mm}$

Data collection

Bruker SMART APEX CCD area-detector diffractometer	10874 measured reflections
Radiation source: fine-focus sealed tube	1955 independent reflections
Graphite monochromator	1385 reflections with $I > 2\sigma(I)$
$\omega$ scan	$R_{\text{int}} = 0.055$
Absorption correction: multi-scan (SADABS; Bruker, 2000)	$\theta_{\text{max}} = 25.5^\circ$ , $\theta_{\text{min}} = 2.1^\circ$
$T_{\text{min}} = 0.957$ , $T_{\text{max}} = 0.997$	$h = -10 \rightarrow 10$
	$k = -12 \rightarrow 13$
	$l = -23 \rightarrow 23$

Refinement

Refinement on $F^2$	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.041$	H-atom parameters constrained
$wR(F^2) = 0.101$	$w = 1/[\sigma^2(F_o^2) + (0.0419P)^2 + 0.1849P]$
$S = 1.04$	where $P = (F_o^2 + 2F_c^2)/3$
1955 reflections	$(\Delta/\sigma)_{\text{max}} < 0.001$
228 parameters	$\Delta\rho_{\text{max}} = 0.14 \text{ e \AA}^{-3}$
0 restraints	$\Delta\rho_{\text{min}} = -0.13 \text{ e \AA}^{-3}$
Primary atom site location: structure-invariant direct methods	

Special details

**Geometry.** All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

**Refinement.** Refinement of  $F^2$  against ALL reflections. The weighted  $R$ -factor  $wR$  and goodness of fit  $S$  are based on  $F^2$ , conventional  $R$ -factors  $R$  are based on  $F$ , with  $F$  set to zero for negative  $F^2$ . The threshold expression of  $F^2 > \sigma(F^2)$  is used only for calculating  $R$ -factors(gt) etc. and is not relevant to the choice of reflections for refinement.  $R$ -factors based on  $F^2$  are statistically about twice as large as those based on  $F$ , and  $R$ -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters ( $\text{\AA}^2$ )

	$x$	$y$	$z$	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.2992 (3)	0.14972 (19)	0.03121 (11)	0.0530 (6)
O2	0.2744 (3)	0.3535 (2)	0.01341 (12)	0.0647 (7)
O3	-0.1643 (3)	-0.2894 (3)	0.23668 (15)	0.0871 (9)
H3A	-0.1272	-0.2265	0.2196	0.131*
N1	0.0397 (3)	-0.2580 (2)	0.13093 (14)	0.0542 (7)

C1	0.3452 (4)	0.2701 (3)	0.03985 (17)	0.0491 (8)
C2	0.4849 (4)	0.2769 (3)	0.08582 (16)	0.0501 (8)
H2A	0.5785	0.2805	0.0576	0.060*
C3	0.4799 (4)	0.1507 (3)	0.12056 (15)	0.0450 (8)
H3B	0.4037	0.1550	0.1574	0.054*
C4	0.6257 (4)	0.0914 (3)	0.14781 (18)	0.0553 (9)
H4A	0.6661	0.1403	0.1852	0.066*
H4B	0.7043	0.0881	0.1126	0.066*
C5	0.5871 (4)	-0.0399 (3)	0.17203 (17)	0.0569 (9)
H5A	0.5236	-0.0338	0.2125	0.068*
H5B	0.6832	-0.0812	0.1845	0.068*
C6	0.4998 (4)	-0.1229 (3)	0.11905 (16)	0.0501 (8)
C7	0.4574 (4)	-0.2407 (3)	0.15386 (18)	0.0585 (10)
H7A	0.5362	-0.2855	0.1750	0.070*
C8	0.3150 (4)	-0.2855 (3)	0.15666 (17)	0.0546 (9)
H8A	0.2984	-0.3620	0.1775	0.066*
C9	0.1825 (4)	-0.2191 (3)	0.12821 (17)	0.0476 (8)
C10	0.2116 (4)	-0.0959 (3)	0.09669 (17)	0.0500 (8)
C11	0.3588 (4)	-0.0537 (3)	0.09191 (15)	0.0417 (8)
C12	0.4121 (4)	0.0698 (3)	0.06425 (16)	0.0446 (8)
H12A	0.4955	0.0533	0.0316	0.053*
C13	0.4820 (5)	0.3915 (3)	0.1308 (2)	0.0783 (12)
H13A	0.4735	0.4647	0.1031	0.118*
H13B	0.5763	0.3953	0.1568	0.118*
H13C	0.3945	0.3870	0.1609	0.118*
C14	0.0693 (4)	-0.0278 (3)	0.0730 (2)	0.0805 (14)
H14A	0.0889	0.0084	0.0294	0.121*
H14B	0.0440	0.0367	0.1048	0.121*
H14C	-0.0161	-0.0850	0.0697	0.121*
C15	0.6148 (4)	-0.1591 (3)	0.0611 (2)	0.0707 (11)
H15A	0.5614	-0.2090	0.0281	0.106*
H15B	0.7000	-0.2056	0.0797	0.106*
H15C	0.6536	-0.0850	0.0399	0.106*
C16	0.0030 (4)	-0.3792 (3)	0.15327 (17)	0.0525 (8)
C17	0.0571 (4)	-0.4863 (3)	0.12167 (19)	0.0626 (10)
H17A	0.1292	-0.4800	0.0867	0.075*
C18	0.0048 (5)	-0.6021 (3)	0.1418 (2)	0.0709 (11)
H18A	0.0405	-0.6733	0.1200	0.085*
C19	-0.0998 (5)	-0.6117 (4)	0.1940 (2)	0.0745 (12)
H19A	-0.1337	-0.6898	0.2080	0.089*
C20	-0.1549 (4)	-0.5067 (4)	0.2258 (2)	0.0701 (11)
H20A	-0.2257	-0.5136	0.2612	0.084*
C21	-0.1051 (4)	-0.3920 (3)	0.20502 (18)	0.0567 (9)

Atomic displacement parameters ( $\text{\AA}^2$ )

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
O1	0.0561 (14)	0.0463 (13)	0.0566 (14)	-0.0012 (12)	-0.0146 (12)	0.0084 (10)
O2	0.0708 (16)	0.0514 (14)	0.0718 (16)	0.0061 (14)	-0.0142 (14)	0.0115 (12)
O3	0.0668 (18)	0.081 (2)	0.114 (2)	-0.0007 (15)	0.0180 (17)	-0.0079 (17)

N1	0.0475 (17)	0.0515 (17)	0.0638 (19)	-0.0068 (15)	0.0000 (15)	0.0104 (14)
C1	0.050 (2)	0.048 (2)	0.049 (2)	-0.0008 (18)	0.0019 (16)	0.0019 (16)
C2	0.046 (2)	0.0471 (18)	0.057 (2)	-0.0002 (17)	0.0028 (17)	-0.0032 (16)
C3	0.0446 (18)	0.0487 (18)	0.0416 (18)	-0.0018 (16)	-0.0005 (16)	0.0006 (15)
C4	0.053 (2)	0.057 (2)	0.057 (2)	-0.0018 (18)	-0.0101 (18)	0.0053 (17)
C5	0.044 (2)	0.066 (2)	0.061 (2)	-0.0025 (18)	-0.0112 (17)	0.0118 (17)
C6	0.047 (2)	0.0504 (19)	0.053 (2)	0.0012 (17)	-0.0003 (18)	0.0115 (15)
C7	0.054 (2)	0.051 (2)	0.071 (2)	0.0098 (19)	-0.0099 (19)	0.0143 (18)
C8	0.059 (2)	0.0432 (19)	0.061 (2)	0.0028 (18)	-0.0053 (19)	0.0095 (17)
C9	0.049 (2)	0.0442 (18)	0.050 (2)	-0.0004 (17)	-0.0018 (16)	0.0050 (15)
C10	0.048 (2)	0.0449 (19)	0.057 (2)	-0.0007 (17)	-0.0079 (17)	0.0057 (15)
C11	0.046 (2)	0.0399 (18)	0.0392 (18)	-0.0007 (15)	-0.0039 (15)	0.0024 (14)
C12	0.0452 (18)	0.0470 (19)	0.0415 (19)	0.0053 (16)	-0.0012 (15)	0.0054 (14)
C13	0.082 (3)	0.061 (2)	0.093 (3)	0.006 (2)	-0.027 (3)	-0.021 (2)
C14	0.052 (2)	0.060 (2)	0.129 (4)	-0.003 (2)	-0.016 (2)	0.033 (2)
C15	0.061 (2)	0.066 (2)	0.085 (3)	0.015 (2)	0.009 (2)	0.004 (2)
C16	0.0445 (19)	0.053 (2)	0.060 (2)	-0.0052 (17)	-0.0041 (18)	0.0100 (17)
C17	0.066 (2)	0.058 (2)	0.064 (2)	-0.006 (2)	0.0036 (19)	0.0054 (19)
C18	0.073 (3)	0.057 (2)	0.083 (3)	-0.011 (2)	-0.002 (3)	0.001 (2)
C19	0.064 (3)	0.064 (3)	0.095 (3)	-0.017 (2)	-0.004 (2)	0.022 (2)
C20	0.051 (2)	0.083 (3)	0.076 (3)	-0.012 (2)	0.007 (2)	0.019 (2)
C21	0.0457 (19)	0.059 (2)	0.065 (2)	-0.0004 (19)	0.0006 (19)	0.0023 (19)

*Geometric parameters (Å, °)*

O1—C1	1.363 (4)	C8—H8A	0.9300
O1—C12	1.451 (3)	C9—C10	1.484 (4)
O2—C1	1.203 (4)	C10—C11	1.348 (4)
O3—C21	1.366 (4)	C10—C14	1.500 (5)
O3—H3A	0.8200	C11—C12	1.506 (4)
N1—C9	1.299 (4)	C12—H12A	0.9800
N1—C16	1.411 (4)	C13—H13A	0.9600
C1—C2	1.507 (5)	C13—H13B	0.9600
C2—C13	1.517 (4)	C13—H13C	0.9600
C2—C3	1.521 (4)	C14—H14A	0.9600
C2—H2A	0.9800	C14—H14B	0.9600
C3—C4	1.506 (4)	C14—H14C	0.9600
C3—C12	1.526 (4)	C15—H15A	0.9600
C3—H3B	0.9800	C15—H15B	0.9600
C4—C5	1.527 (4)	C15—H15C	0.9600
C4—H4A	0.9700	C16—C21	1.388 (5)
C4—H4B	0.9700	C16—C17	1.389 (5)
C5—C6	1.566 (4)	C17—C18	1.382 (5)
C5—H5A	0.9700	C17—H17A	0.9300
C5—H5B	0.9700	C18—C19	1.372 (5)
C6—C7	1.486 (4)	C18—H18A	0.9300
C6—C11	1.520 (4)	C19—C20	1.375 (5)
C6—C15	1.561 (5)	C19—H19A	0.9300
C7—C8	1.317 (4)	C20—C21	1.368 (5)
C7—H7A	0.9300	C20—H20A	0.9300

C8—C9	1.457 (5)		
C1—O1—C12	108.1 (2)	C9—C10—C14	115.3 (3)
C21—O3—H3A	109.5	C10—C11—C12	127.4 (3)
C9—N1—C16	121.4 (3)	C10—C11—C6	124.1 (3)
O2—C1—O1	120.4 (3)	C12—C11—C6	108.4 (3)
O2—C1—C2	128.9 (3)	O1—C12—C11	118.7 (3)
O1—C1—C2	110.7 (3)	O1—C12—C3	104.2 (2)
C1—C2—C13	112.3 (3)	C11—C12—C3	110.7 (2)
C1—C2—C3	101.8 (3)	O1—C12—H12A	107.6
C13—C2—C3	117.4 (3)	C11—C12—H12A	107.6
C1—C2—H2A	108.3	C3—C12—H12A	107.6
C13—C2—H2A	108.3	C2—C13—H13A	109.5
C3—C2—H2A	108.3	C2—C13—H13B	109.5
C4—C3—C2	121.0 (3)	H13A—C13—H13B	109.5
C4—C3—C12	109.7 (3)	C2—C13—H13C	109.5
C2—C3—C12	101.0 (2)	H13A—C13—H13C	109.5
C4—C3—H3B	108.2	H13B—C13—H13C	109.5
C2—C3—H3B	108.2	C10—C14—H14A	109.5
C12—C3—H3B	108.2	C10—C14—H14B	109.5
C3—C4—C5	108.8 (3)	H14A—C14—H14B	109.5
C3—C4—H4A	109.9	C10—C14—H14C	109.5
C5—C4—H4A	109.9	H14A—C14—H14C	109.5
C3—C4—H4B	109.9	H14B—C14—H14C	109.5
C5—C4—H4B	109.9	C6—C15—H15A	109.5
H4A—C4—H4B	108.3	C6—C15—H15B	109.5
C4—C5—C6	115.0 (3)	H15A—C15—H15B	109.5
C4—C5—H5A	108.5	C6—C15—H15C	109.5
C6—C5—H5A	108.5	H15A—C15—H15C	109.5
C4—C5—H5B	108.5	H15B—C15—H15C	109.5
C6—C5—H5B	108.5	C21—C16—C17	118.2 (3)
H5A—C5—H5B	107.5	C21—C16—N1	118.1 (3)
C7—C6—C11	112.6 (3)	C17—C16—N1	123.4 (3)
C7—C6—C15	106.4 (3)	C18—C17—C16	120.5 (4)
C11—C6—C15	111.7 (3)	C18—C17—H17A	119.7
C7—C6—C5	107.1 (3)	C16—C17—H17A	119.7
C11—C6—C5	109.8 (3)	C19—C18—C17	119.8 (4)
C15—C6—C5	109.1 (3)	C19—C18—H18A	120.1
C8—C7—C6	124.0 (3)	C17—C18—H18A	120.1
C8—C7—H7A	118.0	C18—C19—C20	120.4 (4)
C6—C7—H7A	118.0	C18—C19—H19A	119.8
C7—C8—C9	122.1 (3)	C20—C19—H19A	119.8
C7—C8—H8A	118.9	C21—C20—C19	119.6 (3)
C9—C8—H8A	118.9	C21—C20—H20A	120.2
N1—C9—C8	124.5 (3)	C19—C20—H20A	120.2
N1—C9—C10	117.6 (3)	O3—C21—C20	118.3 (3)
C8—C9—C10	117.8 (3)	O3—C21—C16	120.4 (3)
C11—C10—C9	119.2 (3)	C20—C21—C16	121.3 (3)
C11—C10—C14	125.5 (3)		

C12—O1—C1—O2	175.7 (3)	C14—C10—C11—C6	177.4 (3)
C12—O1—C1—C2	-5.8 (3)	C7—C6—C11—C10	-2.1 (5)
O2—C1—C2—C13	34.0 (5)	C15—C6—C11—C10	117.5 (4)
O1—C1—C2—C13	-144.4 (3)	C5—C6—C11—C10	-121.3 (3)
O2—C1—C2—C3	160.4 (3)	C7—C6—C11—C12	174.2 (3)
O1—C1—C2—C3	-18.0 (3)	C15—C6—C11—C12	-66.2 (3)
C1—C2—C3—C4	153.7 (3)	C5—C6—C11—C12	54.9 (3)
C13—C2—C3—C4	-83.3 (4)	C1—O1—C12—C11	151.1 (3)
C1—C2—C3—C12	32.6 (3)	C1—O1—C12—C3	27.3 (3)
C13—C2—C3—C12	155.6 (3)	C10—C11—C12—O1	-8.0 (5)
C2—C3—C4—C5	-173.5 (3)	C6—C11—C12—O1	175.9 (3)
C12—C3—C4—C5	-56.6 (3)	C10—C11—C12—C3	112.5 (4)
C3—C4—C5—C6	51.6 (4)	C6—C11—C12—C3	-63.6 (3)
C4—C5—C6—C7	-173.9 (3)	C4—C3—C12—O1	-166.0 (2)
C4—C5—C6—C11	-51.4 (4)	C2—C3—C12—O1	-37.2 (3)
C4—C5—C6—C15	71.3 (4)	C4—C3—C12—C11	65.3 (3)
C11—C6—C7—C8	4.8 (5)	C2—C3—C12—C11	-165.9 (3)
C15—C6—C7—C8	-117.8 (4)	C9—N1—C16—C21	126.8 (4)
C5—C6—C7—C8	125.6 (4)	C9—N1—C16—C17	-60.1 (5)
C6—C7—C8—C9	-3.2 (6)	C21—C16—C17—C18	-0.4 (5)
C16—N1—C9—C8	-9.8 (5)	N1—C16—C17—C18	-173.5 (3)
C16—N1—C9—C10	173.1 (3)	C16—C17—C18—C19	-0.9 (6)
C7—C8—C9—N1	-178.5 (4)	C17—C18—C19—C20	1.0 (6)
C7—C8—C9—C10	-1.4 (5)	C18—C19—C20—C21	0.1 (6)
N1—C9—C10—C11	-178.8 (3)	C19—C20—C21—O3	178.7 (4)
C8—C9—C10—C11	3.9 (5)	C19—C20—C21—C16	-1.4 (6)
N1—C9—C10—C14	1.7 (5)	C17—C16—C21—O3	-178.6 (3)
C8—C9—C10—C14	-175.6 (3)	N1—C16—C21—O3	-5.1 (5)
C9—C10—C11—C12	-177.6 (3)	C17—C16—C21—C20	1.5 (5)
C14—C10—C11—C12	1.9 (6)	N1—C16—C21—C20	175.0 (3)
C9—C10—C11—C6	-2.0 (5)		

Hydrogen-bond geometry ( $\text{\AA}$ ,  $^\circ$ )

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O3—H3A $\cdots$ N1	0.82	2.28	2.747 (4)	116